

MLLNVLRICI	IVCLVNDGAG	KHSEGRERTK	TYSLNSRGYF	40
RKERGARRSK	ILLVNTKGLD	EPHIGHGDFG	LVAELFDSTR	80
THTNRKEPDM	NKVKLFSTVA	HGNKSARRKA	YNGSRRNIFS	120
RRSFDKRNTE	VTEKPGAKMF	WNNFLVKMNG	APQNTSHGSK	160
AQEIMKEACK	TLPFTQNIVH	ENCDRMVIQN	NLCFGKCISL	200
HVPNQDRRN	TCSHCLPSKF	TLNHLTLNCT	GSKNVVKVVM	240
MVEECTCEAH	KSNFHQTAQF	NMDTSTTLHH		270

Figure 1. Deduced amino acid sequence of *Xenopus cerberus* protein. SEQ ID NO:1.

000001-031101

Figure 2. Nucleotide sequence of the full-length cerberus DNA derived from the *Xenopus* organizer. The sense strand is on top (in the 5' to 3' direction) and the antisense strand on the bottom line (on the opposite direction). SEQ ID NO:2.

GAATCCCAG	CAAGTCGCTC	AGAAACACTG	CAGGGTCTAG	ATATCATACA	ATGTTACTAA	60
CTTAAGGGTC	GTTTCAGCGAG	TCTTTGTGAC	GTCCCAGATC	TATAGTATGT	TACAATGATT	
ATGTACTCAG	GATCTGTATT	ATCGTCTGCC	TTGTGAATGA	TGGAGCAGGA	AAACACTCAG	120
TACATGAGTC	CTAGACATAA	TAGCAGACGG	AACACTTACT	ACCTCGTCCT	TTTGTGAGTC	
AAGGACGAGA	AAGGACAAAA	ACATATTCAC	TTAACAGCAG	AGGTTACTTC	AGAAAAGAAA	180
TTCTGCTCT	TTCTGTTTT	TGTATAAGTG	AATTGTCGTC	TCCAATGAAG	TCTTTTCTTT	
GAGGAGCAG	TAGGAGCAAG	ATTCTGCTGG	TGAATACTAA	AGGTCTTGAT	GAACCCACAC	240
CTCCTCGTC	ATCCTCGTTC	TAAGACGACC	ACTTATGATT	TCCAGAATA	CTTGGGGTGT	
TTGGGCATGG	TGATTTTCGC	TTAGTAGCTG	AACTATTTGA	TTCCACCAGA	ACACATACAA	300
AACCCGTACC	ACTAAAAGCG	AATCATCGAC	TTGATAAACT	AAGTGCGTCT	TGTGTATGTT	
ACAGAAAAGA	GCCAGACATG	AACAAAGTCA	AGCTTTTCTC	AACAGTTGCC	CATGGAACA	360
TGTCTTTTCT	CGGTCTGTAC	TTGTTTCAGT	TCGAAAAGAG	TTGTCAACGG	GTACCTTTGT	
AAAGTGCAAG	AAGAAAAGCT	TACAATGGTT	CTAGAAGGAA	TATTTTTCCT	CGCCGTTCTT	420
TTTCACGTT	TTCTTTTCGA	ATGTTACCAA	GATCTTCCTT	ATAAAAAGGA	GCGGCAAGAA	
TTGATAAAG	AAATACAGAG	GTTACTGAAA	AGCCTGGTGC	CAAGATGTT	TGGAACAATT	480
AACTATTTT	TTTATGTCT	CAATGACTTT	TCGGACCACG	GTTCTACAAG	ACCTTGTTAA	
TTTTGGTTAA	AATGAATGGA	GCCCCACAGA	ATACAAGCCA	TGGCAGTAAA	GCACAGGAAA	540
AAAACCAATT	TTACTTACCT	CGGGGTGTCT	TATGTTCCGT	ACCGTCATTT	CGTGTCTTTT	
TAATGAAAGA	AGCTTGCAAA	ACCTTGTTTT	TCCTCAGAA	TATTGTACAT	GAAACTGTG	600
ATTACTTTCT	TCGAACGTTT	TGGAACAAAA	AGTGAGTCTT	ATAACATGTA	CTTTTGACAC	
ACAGGATGGT	GATACAGAAC	AATCTGTGCT	TTGGTAAATG	CATCTCTCTC	CATGTTCCAA	660
TGTCCTACCA	CTATGTCTTG	TTAGACACGA	AACCATTTAC	GTAGAGAGAG	GTACAAGGTT	
ATCAGCAAGA	TCGACGAAAT	ACTTGTTCCC	ATTGCTTGCC	GTCCAAATTT	ACCCTGAACC	720
TAGTCGTTCT	AGCTGCTTTA	TGAACAAGGG	TAACGAACGG	CAGGTTTAAA	TGGGACTTGG	
ACCTGACGCT	GAATTGTACT	GGATCTAAGA	ATGTAGTAAA	GGTTGTCATG	ATGGTAGAGG	780
TGGACTGCGA	CTTAACATGA	CCTAGATTCT	TACATCATTT	CCAACAGTAC	TACCATCTCC	
AATGCACGTG	TGAAGCTCAT	AAGAGCAACT	TCCACCAAAC	TGCACAGTTT	AACATGGATA	840
TTACGTGCAC	ACTTCGAGTA	TTCTCGTTGA	AGGTGGTTTG	ACGTGTCAAA	TTGTACCTAT	
CATCTACTAC	CCTGCACCAT	TAAAGGACTG	CCATACAGTA	TGGAAATGCC	CTTTTGTTGG	900
GTAGATGATG	GGACGTGGTA	ATTCCTGAC	GGTATGTCAT	ACCTTTACGG	GAAAACAACC	
AATATTTGTT	ACATACTATG	CATCTAAAGC	ATTATGTTGC	CTTCTATTTT	ATATAACCAC	960
TTATAAACAA	TGTATGATAC	GTAGATTTTC	TAATACAACG	GAAGATAAAG	TATATTGGTG	
ATGGAATAAG	GATTGTATGA	ATTATAATTA	ACAAATGGCA	TTTTGTGTAA	CATGCAAGAT	1020
TACCTTATTC	CTAACATACT	TAATATTAAT	TGTTTACCGT	AAAACACATT	GTACGTTCTA	

MSRTRKVDL	LLLAIPGL	ALLPNAYCAS	CEPVRIPMCK	SMPWNMTKMP	NHLHHSTQAN	60
AILAIEQFEG	LLTTECSQDL	LFFLCAMYAP	ICTIDFQHEP	IKPCKSV CER	ARAGCEPILI	120
KYRHTWPESL	ACEELPVYDR	GVCISPEAIV	TVEQGTDSMP	DFSMDSNNGN	CGSGREHCKC	180
KPMKATQKTY	LKNYNYVIR	AKVKEVKV KC	HDATAIVEVK	EILKSSLVNI	PKDTVTLTYN	240
SGCLCPQLVA	NEEYIIMGYE	DKERTRLLLV	EGSLAEKWRD	RLAKVKRWD	QKLRRPRKSK	300
DPVAPIPNKN	SNSRQARS					

Figure 3. Deduced amino acid sequence of *Xenopus* frazzled protein. SEQ ID NO:3.

GAATTTCCCTT	TCACACAGGA	CTCCTGGCAG	AGGTGAATGG	TAGCCCCTAT	GGATTTGGTT	60
CTTAAGGGGAA	AGTGTGTCCT	GAGGACCGTC	TCCACTTACC	AATCGGGATA	CCTAAACCAA	
TGTTGATTTT	GACACATGAT	TGATTGCTTT	CAGATAGGAT	TGAAGGACTT	GGATTTTTAT	120
ACAACTAAAA	CTGTGTACTA	ACTAACGAAA	GTCTATCCTA	ACTTCCTGAA	CCTAAAAATA	
CTAATTCTGC	ACTTTTAAAT	TATCTGAGTA	ATTGTTCAAT	TTGTATTGGA	TGGGACTAAA	180
GATTAAGACG	TGAAAATTTA	ATAGACTCAT	TAACAAGTAA	AACATAACCT	ACCCTGATTT	
GATAAACTTA	ACTCCTTGCT	TTTGACTTGC	CCATAAACTA	TAAGGTGGGG	TGAGTTGTAG	240
CTATTTGAAT	TGAGGAACGA	AAACTGAACG	GGTATTTGAT	ATTCCACCCC	ACTCAACATC	
TTGCTTTTAC	ATGTGCCCAG	ATTTTCCCTG	TATTCCCTGT	ATTCCCTCTA	AAGTAAGCCT	300
AACGAAAATG	TACACGGGTC	TAAAAGGGAC	ATAAGGGACA	TAAGGGAGAT	TTCATTCCGA	
ACACATACAG	GTTGGGCAGA	ATAACAATGT	CTCGAACAAG	GAAAGTGGAC	TCATTACTGC	360
TGTGTATGTC	CAACCCGTCT	TATTGTTTACA	GAGCTTGTTT	CTTTCACCTG	AGTAATGACG	
TACTGGCCAT	ACCTGGACTG	GCGCTTCTCT	TATTACCCAA	TGCTTACTGT	GCTTCGTGTG	420
ATGACCGGTA	TGGACCTGAC	CGCGAAGAGA	ATAATGGGTT	ACGAATGACA	CGAAGCACAC	
AGCCTGTGCG	GATCCCCATG	TGCAAATCTA	TGCCATGGAA	CATGACCAAG	ATGCCCAACC	480
TCGGACACGC	CTAGGGGTAC	ACGTTTAGAT	ACGGTACCTT	GTACTGGTTC	TACGGGTTGG	
ATCTCCACCA	CAGCACTCAA	GCCAATGCCA	TCTTGGCAAT	TGAACAGTTT	GAAGGTTTGC	540
TAGAGGTGGT	GTCTGTAGTT	CGGTTACGGT	AGGACCGTTA	ACTTGTCAAA	CTTCCAAACG	
TGACCACTGA	ATGTAGCCAG	GACCTTTTGT	TCTTTCTGTG	TGCCATGTAT	GCCCCCATTT	600
ACTGGTGACT	TACATCGGTC	CTGGAAAACA	AGAAAGACAC	ACGGTACATA	CGGGGGTAAA	
GTACCATCGA	TTTCAGCAT	GAACCAATTA	AGCCTTGCAA	GTCCGTGTGC	GAAAGGGCCA	660
CATGGTAGCT	AAAGGTCGTA	CTTGGTAAAT	TCGGAAACGTT	CAGGCACACG	CTTTCOCGGT	
GGGCCGGCTG	TGAGCCCATT	CTCATAAAGT	ACCGGCACAC	TTGGCCAGAG	AGCCTGGCAT	720
CCCGGCCGAC	ACTCGGGTAA	GAGTATTTCA	TGGCCGTGTG	AACCGGTCTC	TCGGACCGTA	
GTGAAGAGCT	GCCCGTATAT	GACAGAGGAG	TCTGCATCTC	CCCAGAGGCT	ATCGTCACAG	780
CACTTCTCGA	CGGGCATATA	CTGTCTCCTC	AGACGTAGAG	GGGTCTCCGA	TAGCAGTGTC	
TGGAACAAGG	AACAGATTCA	ATGCCAGACT	TCTCCATGGA	TTCAAACAAT	GGAAATTGCG	840
ACCTTGTTCC	TTGTCTAAGT	TACGGTCTGA	AGAGGTACCT	AAGTTTGTTA	CCTTTAACGC	
GAAGCGGCAG	GGAGCACTGT	AAATGCAAGC	CCATGAAGGC	AACCCAAAAG	ACGTATCTCA	900
CTTCGCCGTC	CCTCGTGACA	TTTACGTTCG	GGTACTTCCG	TTGGGTTTTT	TGCATAGAGT	
AGAATAATTA	CAATTATGTA	ATCAGAGCAA	AAGTGAAAGA	GGTGAAAGTG	AAATGCCACG	960
TCTTATTAAT	GTTAATACAT	TAGTCTCGTT	TTCACTTTCT	CCACTTTCAC	TTTACGGTGC	
ACGCAACAGC	AATTGTGGAA	GTAAAGGAGA	TTCTCAAGTC	TTCCCTAGTG	AACATTCTTA	1020
TGCGTTGTGCG	TTAACACCTT	CATTTCTCT	AAGAGTTTCT	AAGGGATCAC	TTGTAAGGAT	

Figure 6. Nucleotide sequence of the full-length PAPC cDNA derived from the *Xenopus* organizer. The sense strand of the DNA is shown in the top line (in the 5' to 3' direction), and the bottom line shows the antisense strand (opposite orientation). SEQ ID NO:6.

GAATTC	CCAG	AGATGA	ACTC	CTGAGAT	TG	TTTAAAT	GA	CTGCAGG	TCT	GGAAGG	ATTC	60
CTTAAGG	GGTC	TCTACT	TGAG	GAACTCT	AAC	AAAATTT	ACT	GACGTCC	AAG	CCTTCCT	TAAG	
ACATTG	CCAC	ACTGTT	TCTA	GGCATGA	AAAA	AACTGCA	AGT	TTCAACT	TTTG	TTTTTGG	TGC	120
TGTAACG	GTG	TGACAA	AGAT	CCGTACT	TTTT	TTGACGT	TCA	AAGTTGA	AAAC	AAAAACC	CACG	
AACTTTG	ATT	CTTCAAG	ATG	CTGCTT	TCTCT	TCAGAGC	CCAT	TCCAATG	CTG	CTGTTGG	GAC	180
TTGAACT	AA	GAAGTT	TCTAC	GACGAAG	AGA	AGTCTCG	GTA	AGGTTAC	GAC	GACAACC	CTG	
TGATGGT	TTTT	ACAAAC	CAGAC	TGTGAA	AATTG	CCCAGTA	CTA	CATAGAT	GAA	GAAGAACC	CCC	240
ACTACCA	AAA	TGTTTGT	CTG	ACACTTT	AAC	GGGTCAT	GAT	GTATCTA	CTT	CTTCTTG	GGG	
CTGGCA	CTGT	AATTGC	AGTG	TTGTCAC	AAC	ACTCCAT	ATT	TAACACT	TACA	GATATAC	CTG	300
GACCGTG	ACA	TTAACGT	CAC	AACAGT	GTTG	TGAGGT	TATAA	ATTGTG	ATGT	CTATATG	GAC	
CAACCA	ATTT	CCGTCTA	ATG	AAGCAAT	TTA	ATAATT	CCCT	TATCGG	AGTC	CGTGAG	AGTG	360
GTTGGT	TAAA	GGCAGAT	TAC	TTCGTTA	AAT	TATTAAG	GGA	ATAGCCT	CAG	GCACTCT	CAC	
ATGGGC	AGCT	GAGCAT	CATG	GAGAGG	ATTG	ACCGGG	AGCA	AATCTG	CAGG	CAGTCC	CTTC	420
TACCCGT	CGA	CTCGTA	GTAC	CTCTCCT	AAC	TGGCCCT	CGT	TTAGAC	GTCC	GTGAGG	GAAG	
ACTGCA	ACCT	GGCTTT	GAT	GTGGTC	AGCT	TTTCCAA	AGG	ACACTT	CAAG	CTTCTGA	ACG	480
TGACGTT	GGA	CCGAAAC	CTA	CACCACT	CGA	AAAGGTT	TCC	TGTGAAG	TTT	GAAGACT	TGC	
TGAAAGT	GGA	GGTGAG	AGAC	ATTAATG	ACC	ATAGCC	CTCA	CTTTCC	CAGT	GAAATA	ATGC	540
ACTTTC	ACCT	CCACTCT	CTG	TAATTAC	TGG	TATCGG	GAGT	GAAAGG	GTC	CTTTATT	ACG	
ATGTGG	AGGT	GTCTGAA	AGT	TCCTCTG	TGG	GCACCAG	GAT	TCCTTT	AGAA	ATTGCA	ATAG	600
TACACCT	CCA	CAGACTT	TCA	AGGAGAC	ACC	CGTGGT	CCTA	AGGAAAT	CTT	TAACGT	TATC	
ATGAAG	ATGT	TGGGTCC	AAC	TCCATCC	AAG	ACTTTC	AGAT	CTCAAAT	AAT	AGCCACT	TCA	660
TACTTCT	AACA	ACCCAGG	TTG	AGGTAGG	TCT	TGAAAGT	CTA	GAGTTT	TATTA	TCGGTGA	AGT	
GCATTG	ATGT	GCTAACC	AAG	GCAGATG	GGG	TGAAAT	ATGC	AGATTT	AGTC	TTAATG	AGAG	720
CGTAACT	AACA	CGATTGG	TCT	CGTCTAC	CCC	ACTTTAT	ACG	TCTAAAT	CAG	AATTACT	CTC	
AACTGG	ACAG	GGAAATC	CAG	CCAACAT	AACA	TAATGG	AGCT	ACTAGCA	ATG	GATGGGG	GTG	780
TTGACCT	GTG	CCTTTAG	GTC	GGTTGT	ATGT	ATTACCT	CGA	TGATCG	TAC	CTACCCC	CAC	
TACCAT	CACT	ATCTGGT	ACT	GCAGTGG	TTA	ACATCCG	AGT	CCTGGAC	TTT	AATGATA	ACA	840
ATGGTAG	TGA	TAGACCA	TGA	CGTCACCA	AT	TGTAGG	CTCA	GGACCTG	AAA	TTACTATT	GT	
GCCAGT	GTG	TGAGAGA	AGC	ACCATTG	CTG	TGGACCT	AGT	AGAGGAT	GCT	CCTCTGG	GAT	900
CGGGTC	ACAA	ACTCTCT	TCG	TGGTAAC	GAC	ACCTGG	ATCA	TCTCCT	ACGA	GGAGACC	CTA	
ACCTTT	TGTT	GGAGTT	ACAT	GCTACTG	ACG	ATGATGA	AGG	AGTGAAT	GGA	GAAATTG	TTT	960
TGGAAA	ACAA	CCTCAAT	GTA	CGATGAC	TGC	TACTACT	TCC	TCACTTA	CTT	CTTTAAC	AAA	
ATGGAT	TCA	CACTTTG	GCA	TCTCAAG	AGG	TACGTC	AGCT	ATTTAAA	AAT	AACTCC	AGAA	1020
TACCTAA	GTG	GTGAAAC	CGT	AGAGTT	TCTCC	ATGCAGT	CGA	TAAATTT	TAA	TTGAGGT	CTT	

ATTAAATCCA CAGACCTACA GTCAAATATT TGAGGGCCCC TGAAACAGCA CATCAGTCAG	3360
TAATTTAGGT GTCTGGATGT CAGTTTATAA ACTCCCGGGG ACTTTGTCGT GTAGTCAGTC	
GACCTAAAGT GGCCTTTTTA CTTTTCAGAG CTCCTGGGTC TGCCCTCTGT GTTAATCAGC	3420
CTGGATTTC ACGGAAAAAT GAAAATCGTC GAGGACCCAG ACGGGAGACA CAATTAGTCG	
CCCTGGTCAA GTCCTGAGTA GGATCATGGC GTTTTATAT GCATCTCACC TACTTTGGAC	3480
GGGACCAATT CAGGACTCAT CCTAGTACCG CAAAAATATA CGTAGAGTGG ATGAAACCTG	
GTGATTTACA CATAATAGGA AACGCTTGGT TTCAGTGAAG TCTGTGTTGT ATATATTCTG	3540
CACTAAATGT GTATTATCCT TTGCGAACCA AAGTCACTTC AGACACAACA TATATAAGAC	
TTATATACAC GCATTTTGTG TTTGTGTATA TATTTCAGT CCATTCAGAT ATGTGTATAT	3600
AATATATGTG CGTAAACAC AAACACATAT ATAAAGTTCA GGTAAGTCTA TACACATATA	
AGTGCAGACC TTGTAAATTA AATATTCTGA TACTTTTCC TCAATAAATA TTAAAT	
TCACGTCTGG AACATTTAAT TTATAAGACT ATGAAAAAGG AGTTATTTAT AAATTTA	

Fig. 6. (Continuation page 4, SEQ ID NO:6).

FOR THE "FTE" 0000

MVCCGPGRML LGWAGLLVLA ALCLLQVPGA QAAACEPVRI PLCKSLPWNM TKMPNHLHHS	60
TQANAILAME QFEGLLGTHC SPDLLFFLCA MYAPICTIDF QHEPIKPCKS VCERARQGCE	120
PILIKYRHSW PESLACDELP VYDRGVCISP EAIVTADGAD FPMDSSTGHC RGASSERCKC	180
KPV RATQKTY FRNNYNYVIR AKVKEVKMKC HDVTAVVEVK EILKASLVNI PRDTVNLYTT	240
SGCLCPPLTV NEEYVIMGYE DEERSRLLLV EGSIAEKWKD RLGKKVKRWD MKLRHLGLGK	300
TDASDSTQNQ KSGRNSNPRP ARS.	

Figure 7. Deduced amino acid sequence of mouse FRZB-1 protein. SEQ ID NO:7.

000001-0740

Figure 8. Nucleotide sequence of the full-length mouse FRZB-1 cDNA. SEQ ID NO:8.

AAGCCTGGGA CCATGGTCTG CTGCGGCCCG GGACGGATGC TGCTAGGATG GGCCGGGTG	60
TTCGGACCCT GGTACCAGAC GACGCCGGGC CCTGCCTACG ACGATCCTAC CCGGCCCAAC	
CTAGTCCTGG CTGCTCTCTG CCTGCTCCAG GTGCCCGGAG CTCAGGCTGC AGCCTGTGAG	120
GATCAGGACC GACGAGAGAC GGACGAGGTC CACGGGCCTC GAGTCCGACG TCGGACACTC	
CCTGTCCGCA TCCCGCTGTG CAAGTCCCTT CCCTGGAACA TGACCAAGAT GCCCAACCAC	180
GGACAGGCGT AGGGCGACAC GTTCAGGGAA GGGACCTTGT ACTGGTTCTA CGGGTTGGTG	
CTGCACCACA GCACCCAGGC TAACGCCATC CTGGCCATGG AACAGTTCGA AGGGCTGCTG	240
GACGTGGTGT CGTGGGTCCG ATTGCGGTAG GACCGGTACC TTGTCAAGCT TCCCAGACGAC	
GGCACCCACT GCAGCCCGGA TCTTCTCTTC TTCCTCTGTG CAATGTACGC ACCCATTTGC	300
CCGTGGGTGA CGTCGGGCCT AGAAGAGAAG AAGGAGACAC GTTACATGCG TGGGTAAACG	
ACCATCGACT TCCAGCACGA GCCCATCAAG CCCTGCAAGT CTGTGTGTGA GCGCGCCCGA	360
TGGTAGCTGA AGGTCGTGCT CGGGTAGTTC GGGACGTTCA GACACACACT CGCGCGGGCT	
CAGGGCTGCG AGCCCATTTCT CATCAAGTAC CGCCACTCGT GGCCGGAAAG CTTGGCCTGC	420
GTCCCGACGC TCGGGTAAGA GTAGTTCATG GCGGTGAGCA CCGGCCTTTC GAACCGGACG	
GACGAGCTGC CGGTGTACGA CCGCGGCGTG TGCATCTCTC CTGAGGCCAT CGTCACCGCG	480
CTGCTCGACG GCCACATGCT GGCGCCGCAC ACGTAGAGAG GACTCCGGTA GCAGTGGCGC	
GACGGAGCGG ATTTTCCTAT GGATTCAAGT ACTGGACACT GCAGAGGGGC AAGCAGCGAA	540
CTGCCTCGCC TAAAAGGATA CCTAAGTTCA TGACCTGTGA CGTCTCCCCG TTCGTGCTT	
CGTTGCAAAT GTAAGCCTGT CAGAGCTACA CAGAAGACCT ATTTCCGGAA CAATTACAAC	600
GCAACGTTTA CATTCGACA GTCTCGATGT GTCTTCTGGA TAAAGGCCTT GTTAATGTTG	
TATGTCATCC GGGCTAAAGT TAAAGAGGTA AAGATGAAAT GTCATGATGT GACCGCCGTT	660
ATACAGTAGG CCCGATTTC AATTCTCCAT TTCTACTTTA CAGTACTACA CTGGCGGCAA	
GTGGAAGTGA AGGAAATTCT AAAGGCATCA CTGGTAAACA TTCCAAGGGA CACCGTCAAT	720
CACCTTCACT TCCTTTAAGA TTTCCGTAGT GACCATTTGT AAGGTTCCCT GTGGCAGTTA	
CTTTATACCA CCTCTGGCTG CCTCTGTCCT CCACCTACTG TCAATGAGGA ATATGTCATC	780
GAAATATGGT GGAGACCGAC GGAGACAGGA GGTGAATGAC AGTTACTCCT TATACAGTAG	
ATGGGCTATG AAGACGAGGA ACGTTCCAGG TTA CTCTTGG TAGAAGGCTC TATAGCTGAG	840
TACCCGATAC TTCTGCTCCT TGCAAGGTCC AATGAGAACC ATCTTCCGAG ATATCGACTC	
AAGTGGAAGG ATCGGCTTGG TAAGAAAGTC AAGCGCTGGG ATATGAAACT CCGACACCTT	900
TTCACCTTCC TAGCCGAACC ATTCTTTCAG TTCGCGACCC TATACTTTGA GGCTGTGGAA	
GGACTGGGTA AAAGTATGATC TAGCGATTCC ACTCAGAATC AGAAGTCTGG CAGGAATCTC	960
CCTGACCCAT TTTGACTACG ATCGCTAAGG TGAGTCTTAG TCTTCAGACC GTCCTTGAGA	

AATCCCCGGC	CAGCACGCAG	CTAAATCCTG	AAATGTAAAA	GGCCACACCC	ACGGACTCCC	1020
TTAGGGGCCG	GTCGTGCGTC	GATTTAGGAC	TTTACATTTT	CCGGTGTGGG	TGCCTGAGGG	
TTCTAAGACT	GGCGCTGGTG	GACTAACAAA	GGAAAACCGC	ACAGTTGTGC	TCGTGACCGA	1080
AAGATTCTGA	CCGCGACCAC	CTGATTGTTT	CCTTTTGGCG	TGTCAACACG	AGCACTGGCT	
TTGTTTACCG	CAGACACCGC	GTGGCTACCG	AAGTTACTTC	CGGTCCCCTT	TCTCCTGCTT	1140
AACAAATGGC	GTCTGTGGCG	CACCGATGGC	TTCAATGAAG	GCCAGGGGAA	AGAGGACGAA	
CTTAATGGCG	TGGGGTTAGA	TCCTTTAATA	TGTTATATAT	TCTGTTTCAT	CAATCACGTG	1200
GAATTACCGC	ACCCCAATCT	AGGAAATTAT	ACAATATATA	AGACAAAAGTA	GTTAGTGCAC	
GGGACTGTTC	TTTTGCAACC	AGAATAGTAA	ATTAAATATG	TTGATGCTAA	GGTTTCTGTA	1260
CCCTGACAAG	AAAACGTTGG	TCTTATCATT	TAATTTATAC	AACTACGATT	CCAAAGACAT	
CTGGACTCCC	TGGGTTTAAT	TTGGTGTTCT	GTACCCTGAT	TGAGAATGCA	ATGTTTCATG	1320
GACCTGAGGG	ACCCAAATTA	AACCACAAGA	CATGGGACTA	ACTCTTACGT	TACAAAGTAC	
TAAAGAGAGA	ATCCTGGTCA	TATCTCAAGA	ACTAGATATT	GCTGTAAGAC	AGCCTCTGCT	1380
ATTTCTCTCT	TAGGACCAGT	ATAGAGTTCT	TGATCTATAA	CGACATTCTG	TCGGAGACGA	
GCTGCGCTTA	TAGTCTTG TG	TTTGTATGCC	TTTGTCCATT	TCCCTCATGC	TGTGAAAGTT	1440
CGACGCGAAT	ATCAGAACAC	AAACATACGG	AAACAGGTAA	AGGGAGTACG	ACACTTTCAA	
ATACATGTTT	ATAAAGGTAG	AACGGCATT	TGAAATCAGA	CACTGCACAA	GCAGAGTAGC	1500
TATGTACAAA	TATTTCCATC	TTGCCGTAAA	ACTTTAGTCT	GTGACGTGTT	CGTCTCATCG	
CCAACACCAG	GAAGCATTTA	TGAGGAAACG	CCACACAGCA	TGACTTATTT	TCAAGATTGG	1560
GGTTGTGGTC	CTTCGTAAAT	ACTCCTTTGC	GGTGTGTCGT	ACTGAATAAA	AGTTCTAACC	
CAGGCAGCAA	AATAAATAGT	GTTGGGAGCC	AAGAAAAGAA	TATTTTGCCT	GGTTAAGGGG	1620
GTCCGTCGTT	TTATTTATCA	CAACCCTCGG	TTCTTTTCTT	ATAAAACGGA	CCAATTCCCC	
CACACTGGAA	TCAGTAGCCC	TTGAGCCATT	AACAGCAGTG	TTCTTCTGGC	AAGTTTTTGA	1680
GTGTGACCTT	AGTCATCGGG	AACTCGGTAA	TTGTGTCAC	AAGAAGACCG	TTCAAAAAC	
TTTGTTTCATA	AATGTATTCA	CGAGCATTAG	AGATGAACTT	ATAACTAGAC	ATCTGTTGTT	1740
AAACAAGTAT	TTACATAAGT	GCTCGTAATC	TCTACTTGAA	TATTGATCTG	TAGACAACAA	
ATCTCTATAG	CTCTGCTTCC	TTCTAAATCA	AACCCATTGT	TGGATGCTCC	CTCTCCATTC	1800
TAGAGATATC	GAGACGAAGG	AAGATTTAGT	TTGGGTAACA	ACCTACGAGG	GAGAGGTAAG	

MVCGSPGGML LLRAGLLALA ALCLLRVPGA RAAACEPVRI PLCKSLPWNM TKMPNHLHHS 60
TQANAILAIE QFEGLLGTHC SPDLLFFLCA MYAPICTIDF QHEPIKPCKS VCERARQGCE 120
PILIKYRHSW PENLACEELP VYDRGVCISP EAIVTADGAD FPMDSNGNC RGASSERCKC 180
KPIRATQKTY FRNNYNYVIR AKVKEIKTKC HDVTAVVEVK EILKSSLVNI PRDTVNLYTS 240
SGCLCPPLNV NEEYIIMGYE DEERSRLLLV EGSIAEKWKD RLGKKVKRWD MKLRHLGLSK 300
SDSSNSDSTQ SQKSGRNSNP RQARN.

Figure 9. Deduced amino acid sequence of human FRZB-1 protein. SEQ ID NO:9.

Figure 10. Nucleotide sequence of the full-length human FRZB-1 cDNA. SEQ ID NO:10.

This sequence was assembled from public ESTs from the Genbank database

(accession numbers: H18848, R63748, W38677, W44760, H38379 and N71244).

GGCGGAGCGG GCCTTTTGGC GTCCACTGCG CGGCTGCACC CTGCCCCATC TGCCGGGATC 60
CCGCCTCGCC CGGAAAACCG CAGGTGACGC GCCGACGTGG GACGGGGTAG ACGGCCCTAG
ATGGTCTGCG GCAGCCCGGG AGGGATGCTG CTGCTGCGGG CCGGGCTGCT TGCCCTGGCT 120
TACCAGACGC CGTCGGGGCCC TCCCTACGAC GACGACGCCC GGCCCCACGA ACGGGACCGA
GCTCTCTGCC TGCTCCGGGT GCCCGGGGCT CGGGCTGCAG CCTGTGAGCC CGTCCGCATC 180
CGAGAGACGG ACGAGGCCCA CGGGCCCCGA GCCCGACGTC GGACACTCGG GCAGGCGTAG
CCCCTGTGCA AGTCCCTGCC CTGGAACATG ACTAAGATGC CCAACCACCT GCACCACAGC 240
GGGGACACGT TCAGGGACGG GACCTTGTA TATTCTACG GGTGGGTGGA CGTGGTGTGC
ACTCAGGCCA ACGCCATCCT GGCCATCGAG CAGTTCGAAG GTCTGCTGGG CACCCACTGC 300
TGAGTCCGGT TCGGGTAGGA CCGGTAGCTC GTCAAGCTTC CAGACGACCC GTGGGTGACG
AGCCCCGATC TGCTCTTCTT CCTCTGTGCC ATGTACGCGC CCATCTGCAC CATTGACTTC 360
TCGGGGCTAG ACGAGAAGAA GGAGACACGG TACATGCGCG GGTAGACGTG GTAAGTGAAG
CAGCACGAGC CCATCAAGCC CTGTAAGTCT GTGTGCGAGC GGGCCCGGCA GGGCTGTGAG 420
GTCGTGCTCG GGTAGTTCGG GACATTGAGA CACACGCTCG CCCGGGCCGT CCCGACACTC
CCCATACTCA TCAAGTACCG CCACTCGTGG CCGGAGAACC TGGCCTGCGA GGAGCTGCCA 480
GGGTATGAGT AGTTCATGGC GGTGAGCACC GGCCTCTTGG ACCGGACGCT CCTCGACGGT
GTGTACGACA GGGGCGTGTG CATCTCTCCC GAGGCCATCG TTAAGTGGGA CGGAGCTGAT 540
CACATGCTGT CCGCGCACAC GTAGAGAGGG CTCCGGTAGC AATGACGCCT GCCTCGACTA
TTTCCTATGG ATTCTAGTAA CGGAACTGT AGAGGGGCAA GCAGTGAACG CTGTAAATGT 600
AAAGGATACC TAAGATCATT GCCTTTGACA TCTCCCCGTT CGTCACTTGC GACATTTACA
AAGCCTATTA GAGCTACACA GAAGACCTAT TTCCGGAACA ATTACAACCTA TGTCATTCCG 660
TTCGGATAAT CTCGATGTGT CTTCTGGATA AAGGCCCTGT TAATGTTGAT ACAGTAAGCC
GCTAAAGTTA AAGAGATAAA GACTAAGTGC CATGATGTGA CTGCAGTAGT GGAGGTGAAG 720
CGATTTCAAT TTCTCTATTT CTGATTCACG GTACTACACT GACGTCATCA CCTCCACTTC
GAGATTCTAA AGTCCTCTCT GGTAAACATT CCACGGGACA CTGTCAACCT CTATACCAGC 780
CTCTAAGATT TCAGGAGAGA CCATTTGTAA GGTGCCCTGT GACAGTTGGA GATATGGTGC
TCTGGCTGCC TCTGCCCTCC ACTTAATGTT AATGAGGAAT ATATCATCAT GGGCTATGAA 840
AGACCGACGG AGACGGGAGG TGAATTACAA TTACTCCTTA TATAGTAGTA CCCGATACTT

GATGAGGAAC CTACTCCTTG	GTTCAGATT CAAGGTCTAA	ACTCTTGGTG TGAGAACCAC	GAAGGCTCTA CTTCCGAGAT	TAGCTGAGAA ATCGACTCTT	GTGGAAGGAT CACCTTCCTA	900
CGACTCGGTA GCTGAGCCAT	AAAAAGTTAA TTTTTCAATT	GCGCTGGGAT CGCGACCCTA	ATGAAGCTTC TACTTCGAAG	GTCATCTTGG CAGTAGAACC	ACTCAGTAAA TGAGTCATTT	960
AGTGATTCTA TCACTAAGAT	GCAATAGTGA CGTTATCACT	TTCCACTCAG AAGGTGAGTC	AGTCAGAAGT TCAGTCTTCA	CTGGCAGGAA GACCGTCCTT	CTCGAACCCC GAGCTTGGGG	1020
CGGCAAGCAC GCCGTTTCGTG	GCAACTAAAT CGTTGATTTA	CCCGAAATAC GGGCTTTATG	AAAAAGTAAC TTTTTCATTG	ACAGTGGACT TGTCACCTGA	TCCTATTAAG AGGATAATTC	1080
ACTTACTTGC TGAATGAACG	ATTGCTGGAC TAACGACCTG	TAGCAAAGGA ATCGTTTCCT	AAATTGCACT TTTAACGTGA	ATTGCACATC TAACGTGTAG	ATATTCTATT TATAAGATAA	1140
GTTTACTATA CAAATGATAT	AAAATCATGT TTTTAGTACA	GATAACTGAT CTATTGACTA	TATTACTTCT ATAATGAAGA	GTTTCTCTTT CAAAGAGAAA	TGGTTTCTGC ACCAAAGACG	1200
TTCTCTCTTC AAGAGAGAAG	TCTCAACCCC AGAGTTGGGG	TTTGTAATGG AAACATTACC	TTTGGGGGCA AAACCCCCGT	GACTCTTAAG CTGAGAATTC	TATATTGTGA ATATAACACT	1260
GTTTTCTATT CAAAGATAA	TCACTAATCA AGTGATTAGT	TGAGAAAAAC ACTCTTTTTG	TGTTCTTTTG ACAAGAAAAAC	CAATAATAAT GTTATTATTA	AAATTAAACA TTTAATTTGT	1320
TGCTGTTACC ACGACAATGG	AGAGCCTCTT TCTCGGAGAA	TGCTGAGTCT ACGACTCAGA	CCAGATGTTA GGTCTACAAT	ATTTACTTTC TAAATGAAAG	TGCACCCCAA ACGTGGGGTT	1380
TTGGGAATGC AACCCTTACG	AATATTGGAT TTATAACCTA	GAAAAGAGAG CTTTTCTCTC	GTTTCTGGTA CAAAGACCAT	TTACAGAAA AAGTGTCTTT	GCTAGATATG CGATCTATAC	1440
CCTTAAAACA GGAATTTTGT	TACTCTGCCG ATGAGACGGC	ATCTAATTAC TAGATTAATG	AGCCTTATTT TCGGAATAAA	TTGTATGCCT AACATACGGA	TTTGGGCATT AAACCCGTAA	1500
CTCCTCATGC GAGGAGTACG	TTAGAAAGTT AATCTTTCAA	CCAAATGTTT GGTTTACAAA	ATAAAGGTAA TATTTCCATT	AATGGCAGTT TTACCGTCAA	TGAAGTCAAA ACTTCAGTTT	1560
TGTCACATAG ACAGTGTATC	GCAAAGCAAT CGTTTCGTTA	CAAGCACCAG GTTTCGTGGT	GAAGTGTTTA CTTCACAAAT	TGAGGAAACA ACTCCTTTGT	ACACCCAAGA TGTGGGTTCT	1620
TGAATTATTT ACTTAATAAA	TTGAGACTGT AACTCTGACA	CAGGAAGTAA GTCCTTCATT	AATAAATAGG TTATTTATCC	AGCTTAAGAA TCGAATTCTT	AGAACATTTT TCTTGTAATA	1680
GCCTGATTGA CGGACTAACT	GAAGCACAAAC CTTCGTGTTG	TGAAACCACT ACTTTGGTCA	AGCCGCTGGG TCGGCGACCC	GTGTTAATGG CACAATTACC	TAGCATTCTT ATCGTAAGAA	1740
CTTTTGGCAA GAAAACCGTT	TACATTTGAT ATGTAAACTA	TTGTTTCATGA AACAAGTACT	ATATATTAAT TATATAATTA	CAGCATTAGA GTCGTAATCT	GAAATGAATT CTTTACTTAA	1800
ATAACTAGAC TATTGATCTG	ATCTGCTGTT TAGACGACAA	ATCACCATAG TAGTGGTATC	TTTTGTTTAA AAAACAAATT	TTTGCTTCCT AAACGAAGGA	TTTAAATAAA AAATTTATTT	1860
CCCATTGGTG GGGTAACCAC	AAAGTCAAAA TTTCAGTTTT	AAAAAAAAAA TTTTTTTTTT	AAA TTT			